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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/521,335

DATE: 03/27/2000
TIME: 16:40:55

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This Raw Listing contains the General Information
Section and up to first 5 pages.

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1  <110> APPLICANT: Oppmann, Birgit
2      Timans, Jacqueline C.
3      Kastelein, Robert A.
4      Bazan, J. Fernando
5  <120> TITLE OF INVENTION: Mammalian Cytokines; Related Reagents and Methods
6  <130> FILE REFERENCE: DX0935K
7  <140> CURRENT APPLICATION NUMBER: US/09/521,335
8  <141> CURRENT FILING DATE: 2000-03-09
9  <160> NUMBER OF SEQ ID NOS: 13
10 <170> SOFTWARE: PatentIn Ver. 2.0
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20 <222> LOCATION: (213)..(806)
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24      gccccagccc catggacctc cgagcagggg actcgtgggg g atg tta gcg tgc ctg 176
25                                     Met Leu Ala Cys Leu
26                                     -15
27      tgc acg gtg ctc tgg cac ctc cct gca gtg cca gct ctc aat cgc aca 224
28      Cys Thr Val Leu Trp His Leu Pro Ala Val Pro Ala Leu Asn Arg Thr
29      -10 -5 -1 1
30      ggg gac cca ggg cct ggc ccc tcc atc cag aaa acc tat gac ctc acc 272
31      Gly Asp Pro Gly Pro Gly Pro Ser Ile Gln Lys Thr Tyr Asp Leu Thr
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33      cgc tac ctg gag cac caa ctc cgc agc ttg gct ggg acc tat ctg aac 320
34      Arg Tyr Leu Glu His Gln Leu Arg Ser Leu Ala Gly Thr Tyr Leu Asn
35      25 30 35
36      tac ctg ggc ccc cct ttc aac gag cca gac ttc aac cct ccc cgc ctg 368
37      Tyr Leu Gly Pro Pro Phe Asn Glu Pro Asp Phe Asn Pro Pro Arg Leu
38      40 45 50
39      ggg gca gag act ctg ccc agg gcc act gtt gac ttg gag gtg tgg cga 416
40      Gly Ala Glu Thr Leu Pro Arg Ala Thr Val Asp Leu Glu Val Trp Arg
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42      agc ctc aat gac aaa ctg cgg ctg acc cag aac tac gag gcc tac agc 464
43      Ser Leu Asn Asp Lys Leu Arg Leu Thr Gln Asn Tyr Glu Ala Tyr Ser
44      70 75 80

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48      gag ctg cgc cgc agc ctg gcc cac ttc tgc acc agc ctc cag ggc ctg      560
49      Glu Leu Arg Arg Ser Leu Ala His Phe Cys Thr Ser Leu Gln Gly Leu
50      105                      110                      115
51      ctg ggc agc att gcg ggc gtc atg gca gct ctg ggc tac cca ctg ccc      608
52      Leu Gly Ser Ile Ala Gly Val Met Ala Ala Leu Gly Tyr Pro Leu Pro
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55      Gln Pro Leu Pro Gly Thr Glu Pro Thr Trp Thr Pro Gly Pro Ala His
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57      agt gac ttc ctc cag aag atg gac gac ttc tgg ctg ctg aag gag ctg      704
58      Ser Asp Phe Leu Gln Lys Met Asp Asp Phe Trp Leu Leu Lys Glu Leu
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60      cag acc tgg ctg tgg cgc tcg gcc aag gac ttc aac cgg ctc aag aag      752
61      Gln Thr Trp Leu Trp Arg Ser Ala Lys Asp Phe Asn Arg Leu Lys Lys
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63      aag atg cag cct cca gca gct gca gtc acc ctg cac ctg ggg gct cat      800
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67      Gly Phe
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69      ttgtgagagc cagccctgta tgccaacacc tgttgagcca ggagacagaa gctgtgagcc 916
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92      Ala Leu Asn Arg Thr Gly Asp Pro Gly Pro Gly Pro Ser Ile Gln Lys
93      -1 1                      5                      10                      15
94      Thr Tyr Asp Leu Thr Arg Tyr Leu Glu His Gln Leu Arg Ser Leu Ala

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98	Asn Pro Pro Arg Leu Gly Ala Glu Thr Leu Pro Arg Ala Thr Val Asp						
99		50		55		60	
100	Leu Glu Val Trp Arg Ser Leu Asn Asp Lys Leu Arg Leu Thr Gln Asn						
101		65		70		75	
102	Tyr Glu Ala Tyr Ser His Leu Leu Cys Tyr Leu Arg Gly Leu Asn Arg						
103		80		85		90	
104	Gln Ala Ala Thr Ala Glu Leu Arg Arg Ser Leu Ala His Phe Cys Thr						
105		100		105		110	
106	Ser Leu Gln Gly Leu Leu Gly Ser Ile Ala Gly Val Met Ala Ala Leu						
107		115		120		125	
108	Gly Tyr Pro Leu Pro Gln Pro Leu Pro Gly Thr Glu Pro Thr Trp Thr						
109		130		135		140	
110	Pro Gly Pro Ala His Ser Asp Phe Leu Gln Lys Met Asp Asp Phe Trp						
111		145		150		155	
112	Leu Leu Lys Glu Leu Gln Thr Trp Leu Trp Arg Ser Ala Lys Asp Phe						
113		160		165		170	
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132	gct ctt aat cgc aca gga gat cca ggc cct ggc ccc tcc atc cag aaa					96	
133	Ala Leu Asn Arg Thr Gly Asp Pro Gly Pro Gly Pro Ser Ile Gln Lys						
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135	acc tat gac ctc acc cgc tac ctg gag cat caa ctc cgc agc tta gct					144	
136	Thr Tyr Asp Leu Thr Arg Tyr Leu Glu His Gln Leu Arg Ser Leu Ala						
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139	Gly Thr Tyr Leu Asn Tyr Leu Gly Pro Pro Phe Asn Glu Pro Asp Phe						
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141	aat cct cct cga ctg ggg gca gaa act ctg ccc agg gcc acg gtc aac					240	
142	Asn Pro Pro Arg Leu Gly Ala Glu Thr Leu Pro Arg Ala Thr Val Asn						
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144	ttg gaa gtg tgg cga agc ctc aat gac agg ctg cgg ctg acc cag aac					288	

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148      Tyr Glu Ala Tyr Ser His Leu Leu Cys Tyr Leu Arg Gly Leu Asn Arg
149          80                      85                      90                      95
150      cag gct gcc aca gct gaa ctc cga cgt agc ctg gcc cac ttc tgt acc      384
151      Gln Ala Ala Thr Ala Glu Leu Arg Arg Ser Leu Ala His Phe Cys Thr
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153      agc ctc cag ggc ctg ctg ggc agc att gca ggt gtc atg gcg acg ctt      432
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156      ggc tac cca ctg ccc cag cct ctg cca ggg act gag cca gcc tgg gcc      480
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158          130                      135                      140
159      cct ggc cct gcc cac agt gac ttc ctc cag aag atg gat gac ttc tgg      528
160      Pro Gly Pro Ala His Ser Asp Phe Leu Gln Lys Met Asp Asp Phe Trp
161          145                      150                      155
162      ctg ctg aag gag ctg cag acc tgg cta tgg cgt tca gcc aag gac ttc      576
163      Leu Leu Lys Glu Leu Gln Thr Trp Leu Trp Arg Ser Ala Lys Asp Phe
164          160                      165                      170                      175
165      aac cgg ctt aag aag aag atg cag cct cca gca gct tca gtc acc ctg      624
166      Asn Arg Leu Lys Lys Lys Met Gln Pro Pro Ala Ala Ser Val Thr Leu
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187          65                      70                      75
188      Tyr Glu Ala Tyr Ser His Leu Leu Cys Tyr Leu Arg Gly Leu Asn Arg
189          80                      85                      90                      95
190      Gln Ala Ala Thr Ala Glu Leu Arg Arg Ser Leu Ala His Phe Cys Thr
191                      100                      105                      110
192      Ser Leu Gln Gly Leu Leu Gly Ser Ile Ala Gly Val Met Ala Thr Leu
193                      115                      120                      125
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195          130          135          140
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198    Leu Leu Lys Glu Leu Gln Thr Trp Leu Trp Arg Ser Ala Lys Asp Phe
199    160          165          170          175
200    Asn Arg Leu Lys Lys Lys Met Gln Pro Pro Ala Ala Ser Val Thr Leu
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203          195
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207 <213> ORGANISM: rodent
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214          35          40          45
215    Ile Lys Asn Gln Leu Ala Gln Leu Asn Gly Ser Ala Asn Ala Leu Phe
216          50          55          60
217    Ile Ser Tyr Tyr Thr Ala Gln Gly Glu Pro Phe Pro Asn Asn Val Glu
218          65          70          75          80
219    Lys Leu Cys Ala Pro Asn Met Thr Asp Phe Pro Ser Phe His Gly Asn
220          85          90          95
221    Gly Thr Glu Lys Thr Lys Leu Val Glu Leu Tyr Arg Met Val Ala Tyr
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223    Leu Ser Ala Ser Leu Thr Asn Ile Thr Arg Asp Gln Lys Val Leu Asn
224          115          120          125
225    Pro Thr Ala Val Ser Leu Gln Val Lys Leu Asn Ala Thr Ile Asp Val
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227    Met Arg Gly Leu Leu Ser Asn Val Leu Cys Arg Leu Cys Asn Lys Tyr
228          145          150          155          160
229    Arg Val Gly His Val Asp Val Pro Pro Val Pro Asp His Ser Asp Lys
230          165          170          175
231    Glu Ala Phe Gln Arg Lys Lys Leu Gly Cys Gln Leu Leu Gly Thr Tyr
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VERIFICATION SUMMARY
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